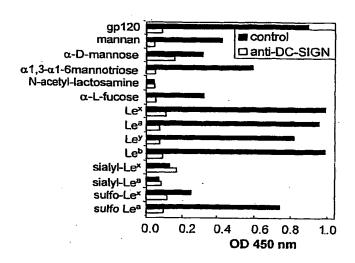
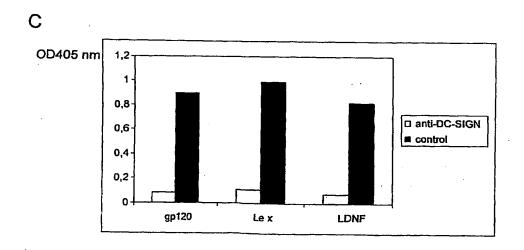
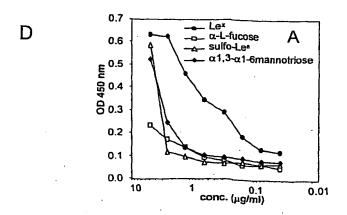
Appl. Serial No. 10/533,981 Applicant: Geijtenbeek et al. Title: "C-type Lectin Binding Molecules, Identification and Uses Thereof" Our Docket: 1943-2 PCT/US Page 1 of 77 REPLACEMENT SHEET

Fig. 1

В

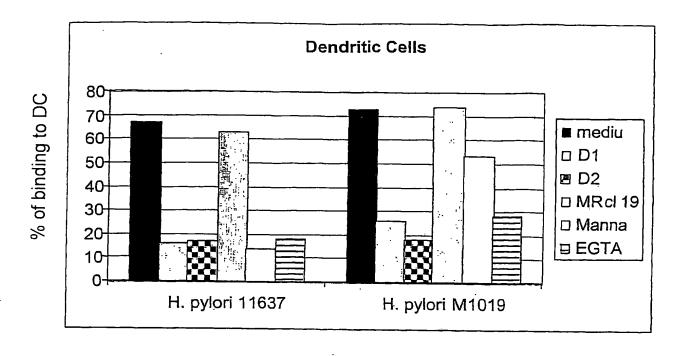






Page 4 of 77 REPLACEMENT SHEET PCT/NL2003/000781

Fig. 4



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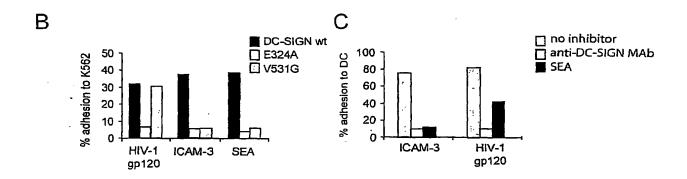
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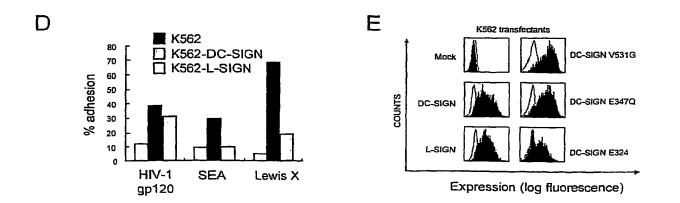
Fig. 19

E324A E347Q V351G

DC-SIGN: WMGLSDLNQEGTWQWVDGSPLLPSFKQYWNRGEPNNVGEEDCAE (SEQ ID NO: 3)
L-SIGN: WMGLSDLNQEGTWQWVDGSPLSPSFQRYWNSGEPNNSGNEDCAE (SEQ ID NO: 4)

Ca²⁺ binding site 1 Ca²⁺ binding site 2 (primary site)





Appl. Serial No. 10/533,981 Applicant: Geijtenbeek et al. Title: "C-type Lectin Binding Molecules, Identification and Uses Thereof" Our Docket: 1943-2 PCT/US Page 44 of 77

REPLACEMENT SHEET

PCT/NL2003/000781

Fig. 44

LPS phase variation in *H. pylori* occurs in vivo.

а	Strain	Number of colonies (%)	Serotype					
		-	Le ^y	mono-Le ^x	(Le ^x) _n	i-antigen	H type 1	
	J223.3	6/30 (20)	+++	-	++	++	+++	
	J223.8	24/30 (80)	-	-	-	+++	+++	

)	Strain	Length of C-tract in gene			Serotype			
		futA	futB	Ley	mono-Le ^x	(Lex) _n	i-antigen	H type 1
	J223.3	n=9 ("off")	n=10 ("on")	+++	-	++	++	+++
	J223.8	n=9 ("off")	n=9 ("off")	~	-	_	, 111	+++
	J223.3 ∆ <i>futB</i>	N.D.	N.D	-	-	-	+++	+++

С	J22	3.3		futA
	1	ATGTTCCAAC	CCCTATTAGA	CGCCTTCATA
	}	GAAAGCGCTT	CCATTGAAAA	AATGGCCTCT
	61	AAATCTCCCC	CCCCCTAA (S	TOP)
	1	. ——	(5	SEQ ID NO: 5)
	J223	3.8		
	1	ATGTTCCAAC	CCCTATTAGA	CGCCTTCATA
		GAAAGCGCTT	CCATTGAAAA	AATGGCCTCT
	61	AAATCTCCCC	CCCCCTAA (STOP)
			(9	SEO ID NO: 5)

J22	3.3 futB
1	ATGTTCCAAC CCCTATTAGA CGCCTTCATA
	GAAAGCGCTT CCATTGAAAA AATGGCCTCT
61	AAATCTCCCC CCCCCTAAA AATC etc.
	(SEQ ID NO: 5)
J22	3.8
1	ATGTTCCAAC CCCTATTAGA CGCCTTCATA
	GAAAGCGCTT CCATTGAAAA AATGGCCTCT
61	AAATCTCCCC CCCCTAA (STOP)
	(SEQ ID NO: 6)